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MAUFFREY, PHILLIPE

PINON-LATAILLADE, GHISLAINE

BIARD, DENIS

ANGULO-MORA, JAINE

<120> SEQUENCES CODING FOR KIN17 PROTEIN AND THEIR APPLICATIONS

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<141> 2000-07-24

OE150> PCT/FR98/02667

<151> 1998-12-09

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Arg Arg Phe Gly Thr Lys Arg Val His Asn Asn Ile Val Tyr Asn

Glu Tyr Ile Ser His Arg Glu His Ile His Met Asn Ala Thr Gln Trp

Glu Thr Leu Thr Asp Phe Thr Lys Trp Leu Gly Arg Glu Gly Leu Cys

Ala Leu Lys Leu Gly Ser Ala Ala Ser Gly Lys Arg Lys Glu Ser

Ser Gln Ser Ser Ala Gln Pro Ala Lys Lys Lys Ser Ala Leu Asp

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Ala Trp Leu Gln Pro Gly Ile Val Val Lys Ile Ile Thr Lys Lys Leu 180 185 190

Gly Glu Lys Tyr His Lys Lys Gly Val Val Lys Glu Val Ile Asp 195 200 205

Arg Tyr Thr Ala Val Val Lys Met Thr Asp Ser Gly Asp Arg Leu Lys 210 225 220

Leu Asp Gln Thr His Leu Glu Thr Val Ile Pro Ala Pro Gly Lys Arg 225 230 235 240

Val Leu Val Leu Asn Gly Gly Tyr Arg Gly Asn Glu Gly Thr Leu Glu 245 250 255

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Ser His Gln Arg Gln Leu Leu Leu Ala Ser Glu Asn Pro Gln Gln Phe 50 55 60

Met Asp Tyr Phe Ser Glu Glu Phe Arg Asn Asp Phe Leu Glu Leu Leu 65 70 75 80

Arg Arg Phe Gly Thr Lys Arg Val His Asn Asn Ile Val Tyr Asn 85 90 95

Glu Tyr Ile Ser His Arg Glu His Ile His Met Asn Ala Thr Gln Trp 100 105 110 Glu Thr Leu Thr Asp Phe Thr Lys Trp Leu Gly Arg Glu Gly Leu Cys 120 Ala Leu Lys Thr Ile Gly Ser Ser Ala Ser Val Lys Arg Lys Glu Ser 130 135 Ser Gln Ser Ser Thr Gln Ser Lys Glu Lys Lys Lys Lys Ser Ala Leu Asp Glu Ile Met Glu Ile Glu Glu Lys Lys Arg Thr Ala Arg Thr Asp Tyr Trp Leu Gln Pro Glu Ile Ile Val Lys Ile Ile Thr Lys Lys Leu Gly Glu Lys Tyr His Lys Lys Lys Ala Ile Val Lys Glu Val Ile Asp Lys Tyr Thr Ala Val Val Lys Met Ile Asp Ser Gly Asp Lys Leu Lys Leu Asp Gln Thr His Leu Glu Thr Val Ile Pro Ala Pro Gly 225 Lys Arg Ile Leu Val Leu Asn Gly Gly Tyr Arg Gly Asn Glu Gly Thr Leu Glu Ser Ile Asn Glu Lys Thr Phe Ser Ala Thr Ile Val Ile Glu Thr Gly Pro Leu Lys Gly Arg Arg Val Glu Gly Ile Gln Tyr Glu Asp 280 Ile Ser Lys Leu Ala 290

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Ser His Gln Arg Gln Leu Leu Leu Ala Ser Glu Asn Pro Gln Gln Phe

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Met Asp Tyr Phe Ser Glu Glu Phe Arg Asn Asp Phe Leu Glu Leu Leu 65 70 75 80

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Glu Tyr Ile Ser His Arg Glu His Ile His Met Asn Ala Thr Gln Trp
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Glu Thr Leu Thr Asp Phe Thr Lys Trp Leu Gly Arg Glu Gly Leu Cys 115 120 125

Lys Val Asp Glu Thr Pro Lys Gly Trp Tyr Ile Gln Tyr Ile Asp Arg 130 135 140

Asp Pro Glu Thr Ile Arg Arg Gln Leu Glu Leu Glu Lys Lys Lys 145 150 155 160

Gln Asp Leu Asp Asp Glu Glu Lys Thr Ala Lys Phe Ile Glu Glu Gln
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Val Arg Arg Gly Leu Glu Gly Lys Glu Gln Glu Thr Pro Val Phe Thr 180 185 190

Glu Leu Ser Arg Glu Asn Glu Glu Glu Lys Val Thr Phe Asn Leu Asn 195 200 205

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Arg Lys Glu Ser Ser Gln Ser Ser Ala Gln Pro Ala Lys Lys Lys Lys 245 250 255

Ser Ala Leu Asp Glu Ile Met Glu Leu Glu Glu Glu Lys Lys Arg Thr 260 265 270

Ala Arg Thr Asp Ala Trp Leu Gln Pro Gly Ile Val Val Lys Ile Ile 275 280 285

Thr Lys Lys Leu Gly Glu Lys Tyr His Lys Lys Lys Gly Val Val Lys 290 295 300

Glu Val Ile Asp Arg Tyr Thr Ala Val Val Lys Met Thr Asp Ser Gly 305 310 315 320

Asp Arg Leu Lys Leu Asp Gln Thr His Leu Glu Thr Val Ile Pro Ala 325 330 335

Pro Gly Lys Arg Val Leu Val Leu Asn Gly Gly Tyr Arg Gly Asn Glu 340 345 350

Gly Thr Leu Glu Ser Ile Asn Glu Lys Ala Phe Ser Ala Thr Ile Val

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Lys Gln Cys Arg Asp Glu Asn Gly Phe Lys Cys His Cys Met Ser Glu 35 40 45

Ser His Gln Arg Gln Leu Leu Leu Ala Ser Glu Asn Pro Gln Gln Phe 50 55 60

Met Asp Tyr Phe Ser Glu Glu Phe Arg Asn Asp Phe Leu Glu Leu Leu 65 70 75 80

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Asp Pro Glu Thr Ile Arg Arg Gln Leu Glu Leu Glu Lys Lys Lys 145 150 155 160

Gln Asp Leu Asp Asp Glu Glu Lys Thr Ala Lys Phe Ile Glu Glu Gln 165 170 175

Val Arg Arg Gly Leu Glu Gly Lys Glu Gln Glu Val Pro Thr Phe Thr 180 185 190

Glu Leu Ser Arg Glu Asn Asp Glu Glu Lys Val Thr Phe Asn Leu Ser 195 200 205

Lys Gly Ala Cys Ser Ser Ser Gly Ala Thr Ser Ser Lys Ser Ser Thr 210 215 220

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280

295

Thr Tyr Arg Gly Leu Ala Ser Pro Ile Leu Glu Ser Glu Arg Leu Tyr

275

305

Ser Leu Glu Ala Leu Ala

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<212> PRT

<213> Mus sp.

<400> 36

Pro Arg Gly Leu Tyr Ile Leu Glu Val Ala Leu Val Ala Leu Leu Tyr

1 10 15

Ser Ile Leu Glu Ile Leu Glu Thr His Arg Leu Tyr Ser Leu Tyr Ser 20 25 30

Leu Glu Gly Leu Tyr Gly Leu Leu Tyr Ser Thr Tyr Arg His Ile Ser 35 40 45

Leu Tyr Ser Leu Tyr Ser Leu Tyr Ser Gly Leu Tyr Val Ala Leu Val 50 55 60

Ala Leu Leu Tyr Ser Gly Leu Val Ala Leu Ile Leu Glu Ala Ser Pro 65 70 75 80

Ala Arg Gly Thr Tyr Arg Thr His Arg Ala Leu Ala Val Ala Leu Val 85 90 95

Ala Leu Leu Tyr Ser Met Glu Thr Thr His Arg Ala Ser Pro Ser Glu 100 105 110

Arg Gly Leu Tyr Ala Ser Pro Ala Arg Gly Leu Glu Leu Tyr Ser Leu
115 120 125

Glu Ala Ser Pro Gly Leu Asn Thr His Arg His Ile Ser Leu Glu Gly 130 135 140

Leu Thr His Arg Val Ala Leu Ile Leu Glu Pro Arg Ala Leu Ala Pro 145 150 155 160

Arg Gly Leu Tyr Leu Tyr Ser Ala Arg Gly Val Ala Leu Leu Glu Val 165 170 175

Ala Leu Leu Glu Ala Ser Asn Gly Leu Tyr Gly Leu Tyr Thr Tyr Arg 180 185 190

Ala Arg Gly Gly Leu Tyr Ala Ser Asn Gly Leu Gly Leu Tyr Thr His 195 200 205

Arg Leu Glu Gly Leu Ser Glu Arg Ile Leu Glu Ala Ser Asn Gly Leu 210 215 220

Leu Tyr Ser Ala Leu Ala Pro His Glu Ser Glu Arg Ala Leu Ala Thr 225 230 235 240

His Arg Ile Leu Glu Val Ala Leu Ile Leu Glu Gly Leu Thr His Arg

245

250

255

Gly Leu Tyr Pro Arg Leu Glu Leu Tyr Ser Gly Leu Tyr Ala Arg Gly 260 265 270

Ala Arg Gly Val Ala Leu Gly Leu Gly Leu Tyr Ile Leu Glu Gly Leu 275 280 285

Asn Thr Tyr Arg Gly Leu Ala Ser Pro Ile Leu Glu Ser Glu Arg Leu
290 295 300

Tyr Ser Leu Glu Ala Leu Ala 305 310